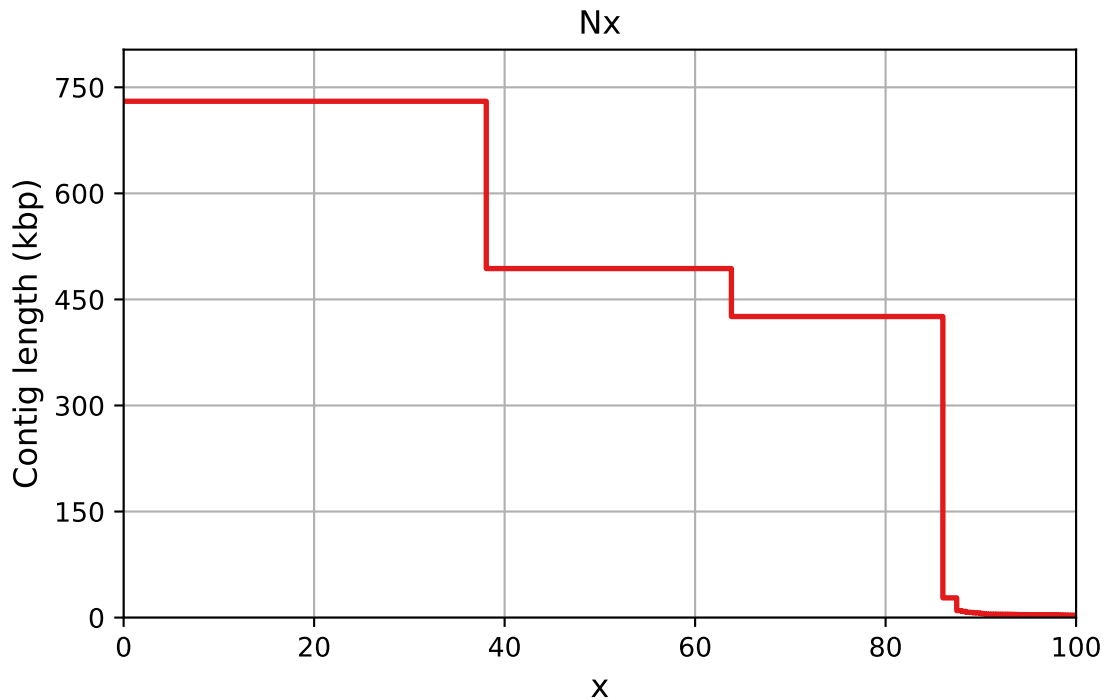


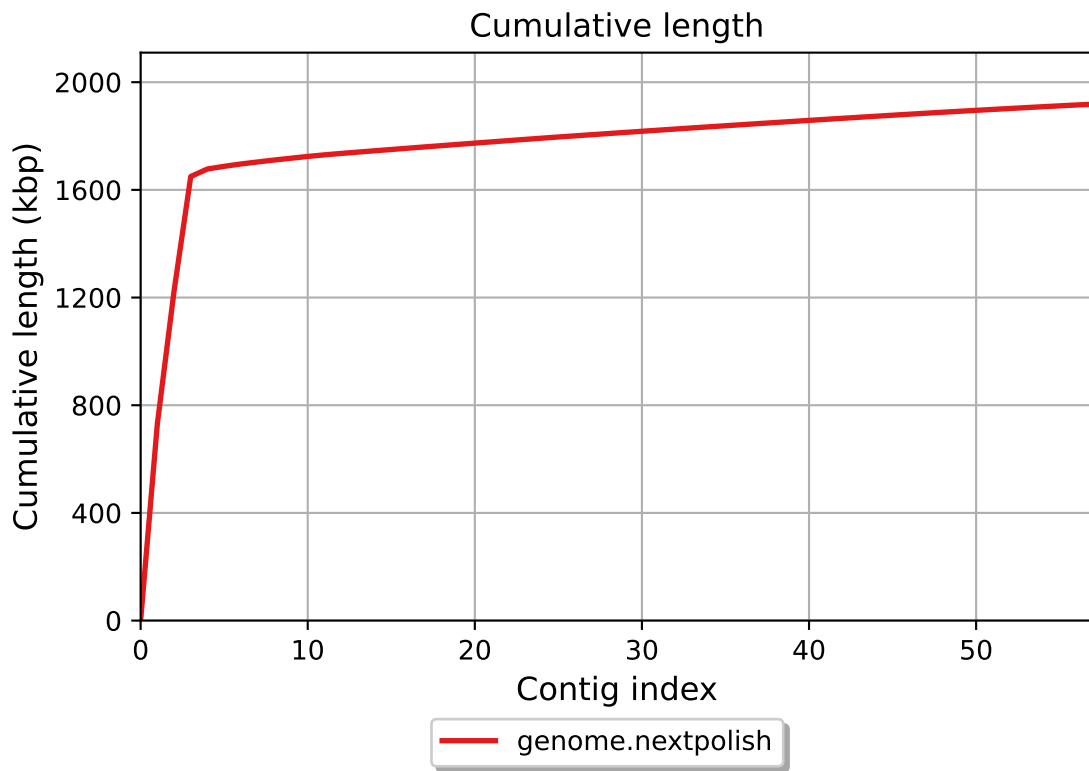
Report

	genome.nextpolish
# contigs (>= 0 bp)	57
# contigs (>= 1000 bp)	57
# contigs (>= 5000 bp)	12
# contigs (>= 10000 bp)	4
# contigs (>= 25000 bp)	4
# contigs (>= 50000 bp)	3
Total length (>= 0 bp)	1918001
Total length (>= 1000 bp)	1918001
Total length (>= 5000 bp)	1735321
Total length (>= 10000 bp)	1677599
Total length (>= 25000 bp)	1677599
Total length (>= 50000 bp)	1649720
# contigs	57
Largest contig	730258
Total length	1918001
GC (%)	31.50
N50	493649
N90	5922
auN	500649.4
L50	2
L90	11
# N's per 100 kbp	0.00
Complete BUSCO (%)	0.00
Partial BUSCO (%)	0.00
# predicted rRNA genes	5 + 5 part

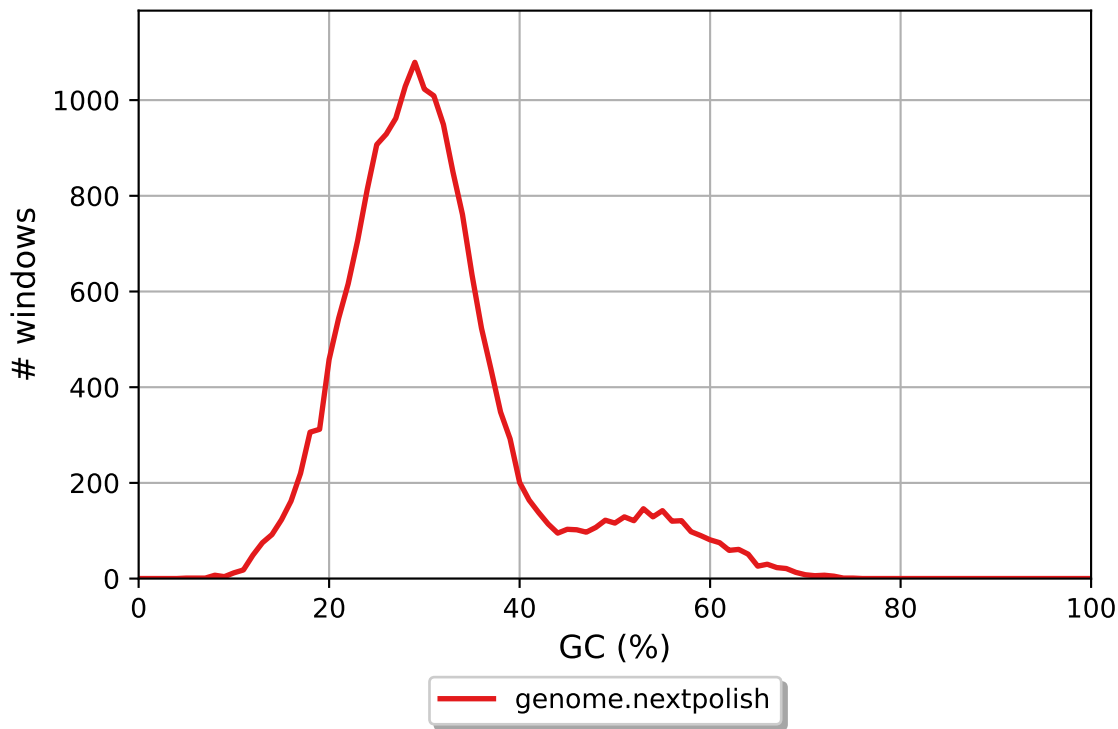
All statistics are based on contigs of size >= 500 bp, unless otherwise noted (e.g., "# contigs (>= 0 bp)" and "Total length (>= 0 bp)" include all contigs).



— genome.nextpolish



GC content



genome.nextpolish GC content

